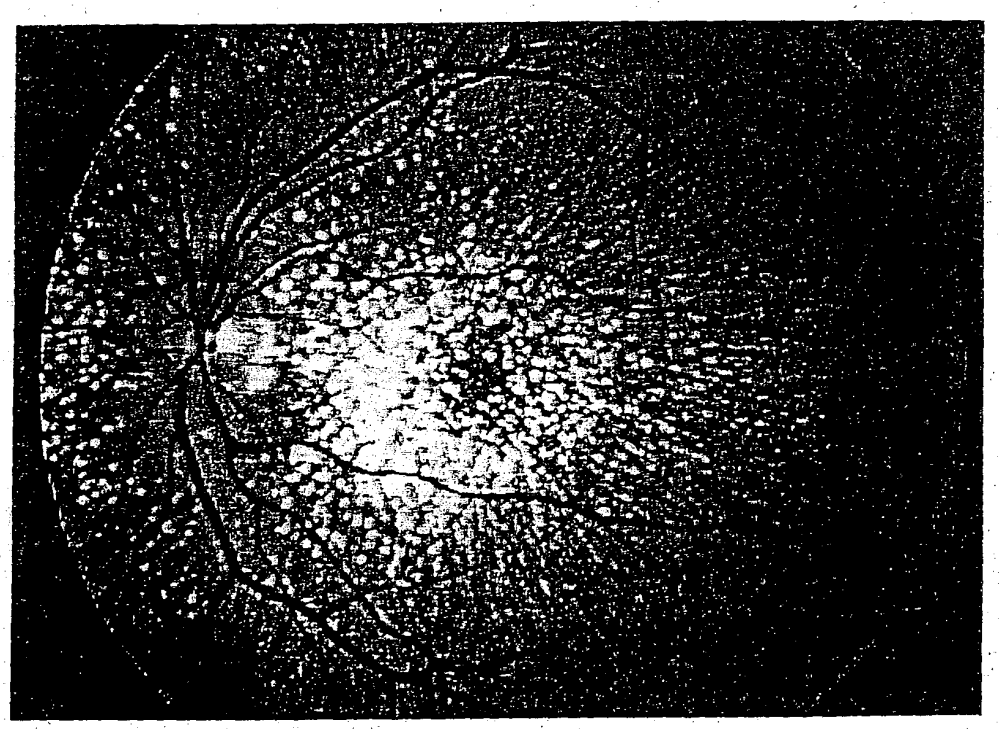

FIG. 1

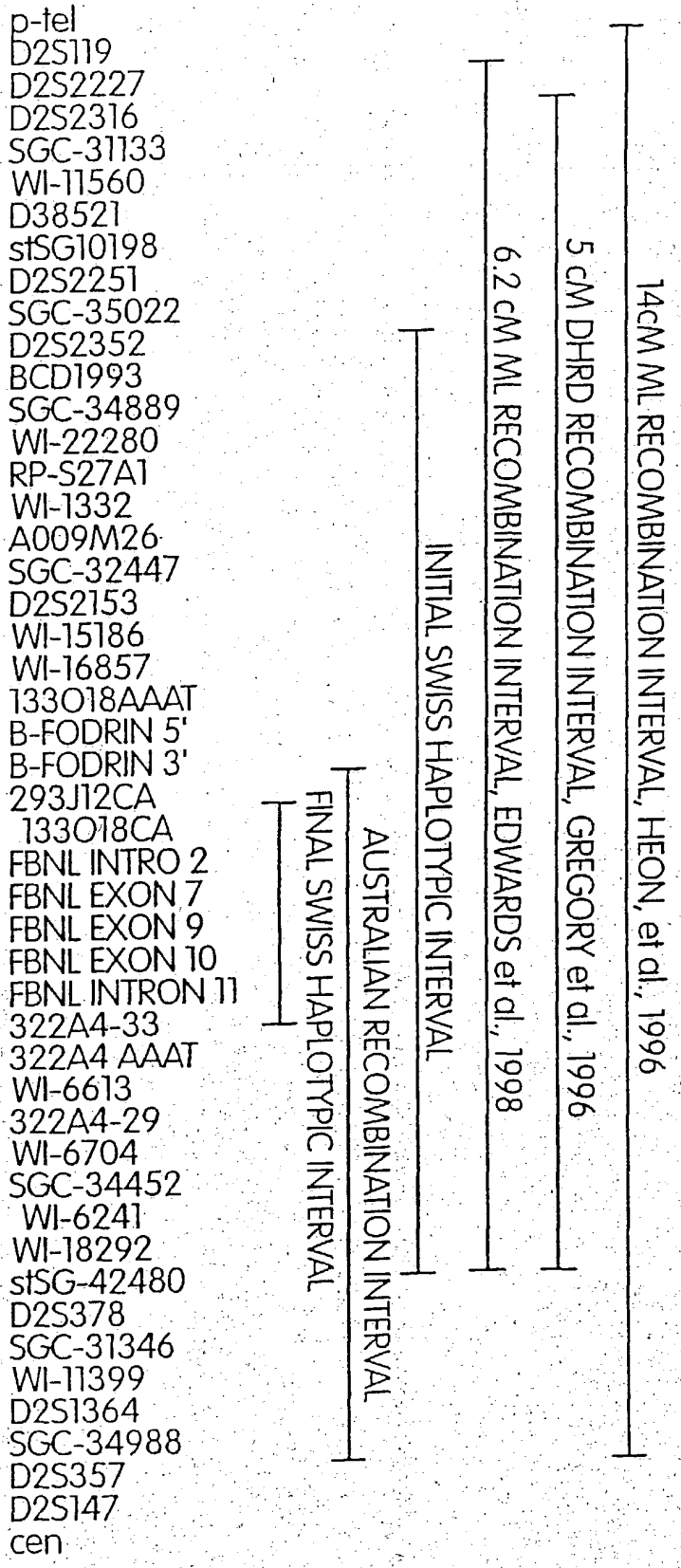
p-tel
D2S119
D2S2227
D2S2316
SGC-31133
WI-11560
D38521
stSG10198
D2S2251
SGC-35022
D2S2352
BCD1993
SGC-34889
WI-22280
RP-S27A1
WI-1332
A009M26
SGC-32447
D2S2153
WI-15186
WI-16857
133O18AAAT
B-FODRIN 5'
B-FODRIN 3'
293J12CA
133O18CA
FBNL INTRO 2
FBNL EXON 7
FBNL EXON 9
FBNL EXON 10
FBNL INTRON 11
322A4-33
322A4 AAAT
WI-6613
322A4-29
WI-6704
SGC-34452
WI-6241
WI-18292
stSG-42480
D2S378
SGC-31346
WI-11399
D2S1364
SGC-34988
D2S357
D2S147
cen
FINAL SWISS HAPLOTYPIC INTERVAL
AUSTRALIAN RECOMBINATION INTERVAL
INITIAL SWISS HAPLOTYPIC INTERVAL
6.2 cM ML RECOMBINATION INTERVAL, EDWARDS et al., 1998
5 cM DHRD RECOMBINATION INTERVAL, GREGORY et al., 1996
14cM ML RECOMBINATION INTERVAL, HEON, et al., 1996

FIG. 2A

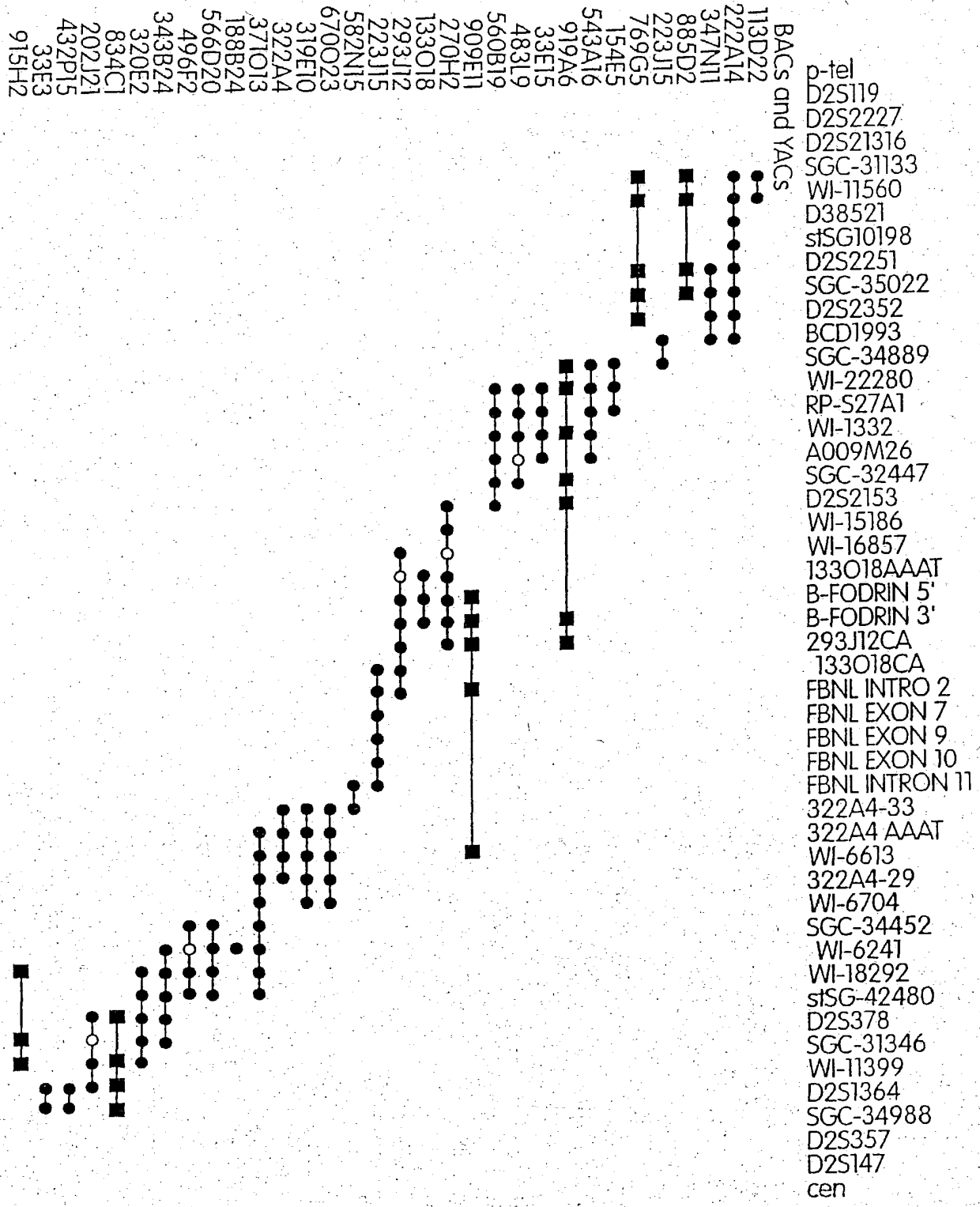
BACs and YACs
113D22
222A14
347N11
885D2
223J15
769G5
154E5
543A16
919A6
33E15
483L9
560B19
909E11
270H2
293J12
133O18
582N15
670O23
319E10
322A4
377O13
188B24
566D20
496F2
343B24
320E2
834C1
202J21
432P15
33E3
915H2
p-tel
D2S119
D2S2227
D2S21316
SGC-31133
WI-11560
D38521
stSG10198
D2S2251
SGC-35022
D2S2352
BCD1993
SGC-34889
WI-22280
RP-S27A1
WI-1332
A009M26
SGC-32447
D2S2153
WI-15186
WI-16857
133O18AAAT
B-FODRIN 5'
B-FODRIN 3'
293J12CA
133O18CA
FBNL INTRO 2
FBNL EXON 7
FBNL EXON 9
FBNL EXON 10
FBNL INTRON 11
322A4-33
322A4 AAAT
WI-6613
322A4-29
WI-6704
SGC-34452
WI-6241
WI-18292
stSG-42480
D2S378
SGC-31346
WI-11399
D2S1364
SGC-34988
D2S357
D2S147
cen

FIG. 2B

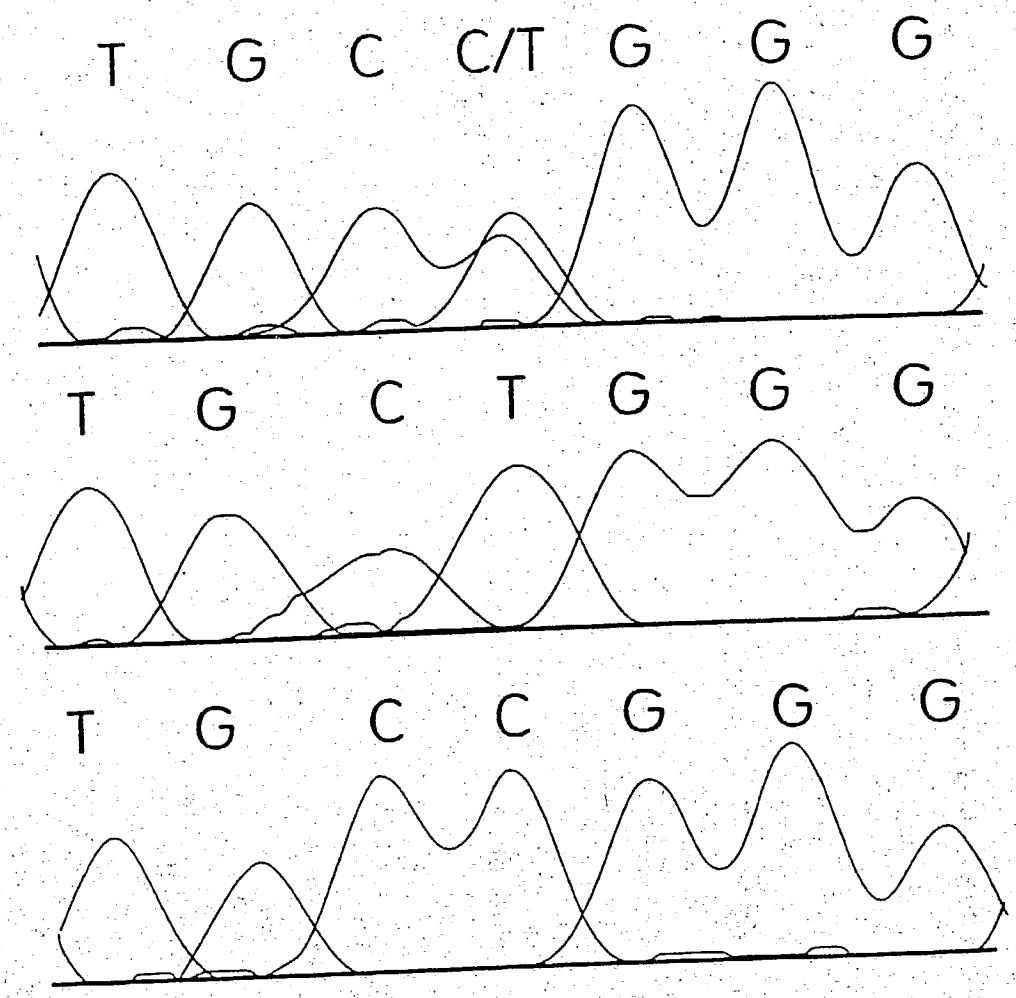

FIG. 3

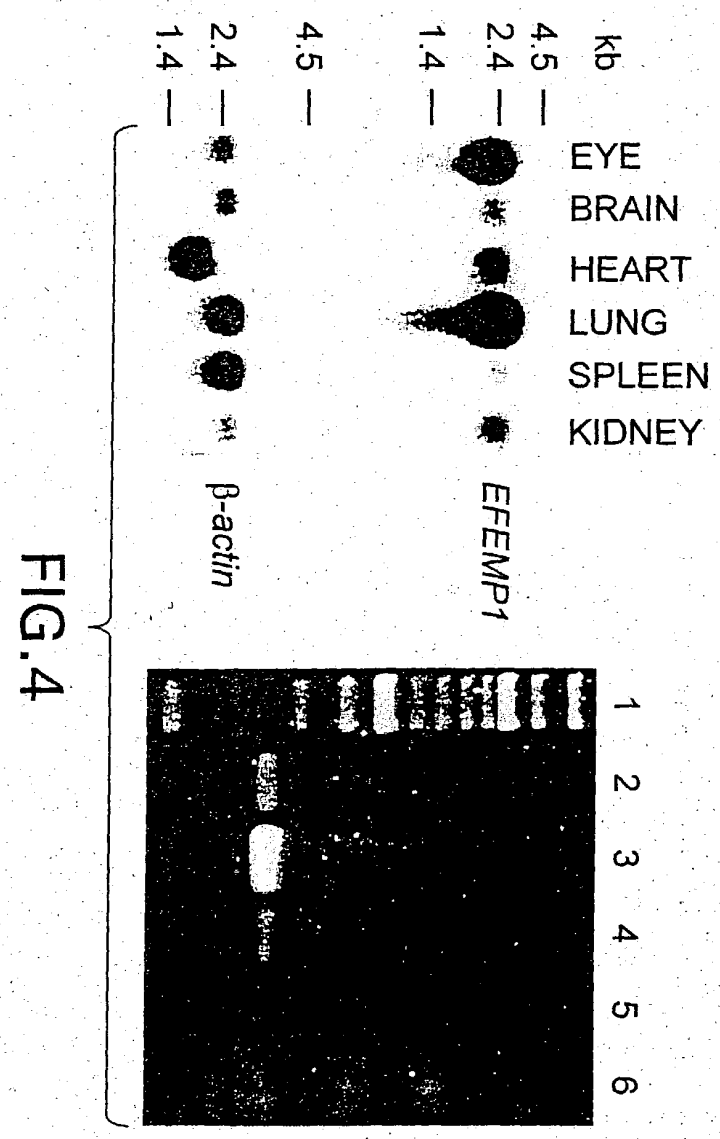
kb
4.5
2.4
1.4
EYE
BRAIN
HEART
LUNG
SPLEEN
KIDNEY
EFEMP1
β-actin
1
2
3
4
5
6

FIG.4

Exon 1

Currently undetermined

exon 2 partial transcript

AGTGTGCGAGATTTAAGCCGCACCTGGATTCCATAGGAGCTGGTTAGAAGCTGGGACGCTGAGCAGCT
CCAGGGGACCGCCGCGTTAGCTTGCTGTTAAGAAAGGGGACCTCATCTCCCTGCCGGGCCAGGCCGCC
CGCCCGAAACTGGTACCTTGGGCTGCGGTGCGATCCCTGGTTCCGGTCCTAGGCAGCCTGAAACCGAA
GGTAGCGTGTCGGGGACCCAGACTGATAAGACAAAAGAGAATCAGTCGCTTTGGGCTGCCCCTCCACA
CAACCTGGGACTTTTAAACAAAGCTGTGCGCAGAGAAAGGCGTGGAAATGCCACTTTGAGAGTTTGTG
CTGGGGGATGTGAGAAGCTCTGAGACATGTGAGAAGGTCTAGTATTCTACTAGAACTGGAAGATTGCT
CTCCGAGTTTTGTTTTGTTATTTTGTTTAAAAAATAAAAAGCTTGAGGCCAAGGCAATTCATATTGGCT
CACAGGTATTTTTGCTGTGCTGTGCAAGGAACTCTGCTAGCTCAAG (SEQ ID No. 46)

Complete intron 2

gtgagtatac tgggaaagcc ttaacgttct agagaacaga gttttgcagccccacctcca
tccccccgggg tgtgggactg gggggcagga aatgcaccgtccccttgaa atggattact
gttttttcct ttcgaccccc tctttctgcagcctgctttg taggtgcagt ataaaatgca
cgctgaatgt cttttgtatgtaaacagcgt agcaggatgg agtaacgtga aatgcaattc
tacagcagtttttacgtctt tgctgcctcg ttcgttggct accgagaagg ttcaggaggggggaggggaga
tgagaaagca gattggaagt tgagtatggt ggtagcctcagcctctccca ccctcctttc
ctcgcttgtg ctcactgcta aagttttgttactttccccg cagcagatac taaacattag
tttgtcctgt attttctttgag (SEQ ID No. 47)

Exon 3

ATTCACAATG TTGAAAGCCC TTTTCCTAAC TATGCTGACT CTGGCGCTGG TCAAGTCACA
GGACACCGAA GAAACCATCA CGTACACG (SEQ ID No. 48)

Start intron 3

gtaaggggtg atggaatttg aaataagtac ggttcagcgg gattctgtgacaaaaaaaga ctttacacac
ccctgccagt gtatttgggc tatactctgctgagggtgat aaattaaaca acacttcatt catgcttcat
atctaagattcgttgtaaat tgccccccttg atcctttcaa aagttcattg ggctcaccacctaagatagg
aaccaacatg taatcatttg tgcagggcta aaaatgggatccgttcaaaa actaaaacca aagaaagtta
catgtttcca aaacattcaacaaattaatg ggtgtaagga actggaaaac ctggactcct
accacatgcagataaaacca atacgtgcag aataagactc aagtcaagta agaacgttaaacaccataaa
gacacatggccttctttgtgtacatgacatgcattctcaagtaagtggcctttattgaatttataaaggctata
tattcattctttttgtataacttg ataattctaataaataaaggcagacaacagtttatgtgttaccaggatg
catattggctaaagtggttttaaaacgtaatgtgtgcaactccgttttgcattttctaattagcgtct
ctgatatttccaagtaatatttgattagttagttgcataggtgtaaccaatgtttaataaaatattaaaaagat
cacctgacccctcccactgctacaaatagttgtggtgagaac agagaaggacagtactgacttcacttctggt
gagtttgtttgcacctctgttctgtgttttcttgtctttaatcagtgttaggcaaatgacatttgtcctggatt
ggaatatgaaaa gcacatttttctactgctcccagtttaaaattaagtaatcctactcgaaagaatgtgaaa
aatttttgaaaagaaaactcttaaaaatgaactaatgtcaattactgataataaac attatctcacttttttgg
taccacattatccctagaatgttagtattcatctggcaaatgttcctttctgttgttcagatccactataaat
aaaatagcttaatacaaatattttgttaact tcatgttaaatgcagttgcttcctctgctgaagataaattaa
gcaagaaaaatgaaggcatgtgctgkttatcttaaaatgaaaatgktttgktattcagactaaacttactgcc
ttctcangggagctaaaattaaattcactacccacttttataatcatctcataaaagattttacttctttttcca
gttgc (SEQ ID No. 49)

FIG.5A

End intron 3

aaaatgcctg gttttcagaa ggggggtycc ataggctgtt tgttcagnggggngctaar gaggnnattt
gaagctgggg actgttnggg ggaaccccnctgttcctttg gagcttaant tanggagcng gtcaggggg
gaaanggaggggctttaatn ctgtnanagg ntttnaaaaa aaaaaaaaaa ntccnggctggttnggggt
gggggnggggg gaaagggcca agaaaaaaaa aaaaaatggtnttttttttt tttaacattt ccaatgtggg
aaaaaaggca aattaataaagagcagtcag agaagttgga gaagattagt ctcaaaacag
aaaagaagatggtactgggc anctgtacca aaaagaacag aagagtttag gcagctgatggttgagaatg
gacccccgaa gctgtccaat gcacagactt gtcttttgaaaaaaaagcga tagaatgtta aaccacccat
ctcatcatat atctaggactttagcacaag gattgttgcc ataagaatga agcttttaga
gtgatttcttagggaatgga cacaccaatt aactgtctcc tggccccacc tttgatgttttcttcacag
(SEQ ID No. 50)

Exon 4

CAATGCACTG ACGGATATGA GTGGGATCCT GTGAGACAGC AATGCAAAG (SEQ ID No. 51)

complete intron 4

gtgagccagcttcaaagacttccatgtgatcattgccttctgtctccatctcttgtgtcactcttcctgtcctgt
ctgtgttataccaaaaaggcatgagcattatatttacatgtttgatttttccctcttagaarattcctgacttat
tttattactgaccacag (SEQ ID No. 52)

Exon 5

ATATTGATGA ATGTGACATT GTCCCAGACG CTTGTAAAGG TGGAATGAAG TGTGTCAACC
ACTATGGAGG ATACCTCTGC CTTCCGAAAA CAGCCCAGAT TATTGTCAAT AATGAACAGC
CTCAGCAGGA AACACAACCA GCAGAAGGAA CCTCAGGGGC AACCACCGGG GTTGTAGCTG
CCAGCAGCAT GGCAACCAGT GGAGTGTTGC CCGGGGGTGG TTTTGTGGCC AGTGCTGCTG
CAGTCGCAGG CCCTGAAATG CAGACTGGCC GAAATAACTT TGTCATCCGG CGGAACCCAG
CTGACCCTCA GCGCATTCCC TCCAACCCTT CCCACCGTAT CCAGTGTGCA GCAGGCTACG
AGCAAAGTGA ACACAACGTG TGCCAAG (SEQ ID No. 53)

Start intron 5

Gtaggaataacttattttcctaacccattgaaagctgcaggagaacccccatgaagcttgaattcggatcca
cgttg (SEQ ID No. 54)

End intron 5

Tttggcattctgaaattttgcaatgaaagatttatacawacatgttgccctgctgtttggcaaaaaacaaacagc
tacmgggraacggtataatattaaaggttgatwacacccagttattggttagatttttagaaatttgtcaatgga
aattatctcaaatacaatatattggatggaaaagcaagtatcatacaatctattaaaatttttaacatacaaaac
aataccatatgttctaatggatgcatccctgtctaacaaaagtacaaaaacatctcagggaaggattcattccta
ccgagacagtggtagctgatgggtcaagggatgaggatggtgtgaggctttagctgtatctgaaatgtttcttaa
caaaacaaaatgagccaagaccaacatgacaaaatgttagcatttgttaaatctgagcagtactcactggtattt
gcaaaattattttctgaacacttgaaataatttataatttaaacatttccaatgcaagaacattataaacttttt
aagaataaagtwaaaatttagcttaagaagtggcmaaatggargaaatatcaacatcttcacaactgacaatyyt
tytstgtgctwgyatgtctswgacag (SEQ ID No. 55)

Exon 6

ACATAGACGA GTGCACTGCA GGGACGCACA ACTGTAGAGC AGACCAAGTG TGCATCAATT
TACGGGGATC CTTTGCATGT CAGTGCCCTC CTGGATATCA GAAGCGAGGG GAGCAGTGCG TAG
(SEQ ID No. 56)

FIG.5B

Start intron 6

Gtaagtaccacagtgcagtgagcacggtcctgtcttgagtaccttaactgtttcaacctcaagcattccaatc
aaagcattcatgtttctttggagagtggtagccaataattccttatttttttatagactaccaatccatttc
cacaataacaagaaacaaccttaaaggttgaggcaggagaaccccatgaagcttgaattc
(SEQ ID No. 57)

End intron 6

Aaactgacatcttatatatatatgcaatagtctcaattattttgttttctttaaatctcaatccacccaccaa
gtttatttaccactgaatggcatgaacattgagtctttgttcttaacttcttaactcagaatacaaagtatat
ttaaaatacatataccctaatttttaacaaaataggaaattattacttttaaaaagagattttctctacatagg
ttttctagataatgcttttcagagaatgctaattcaataatttggttctctttgtgtgtgtgcctgataacctag
(SEQ ID No. 58)

Exon 7

ACATAGATGA ATGTACCATC CCTCCATATT GCCACCAAAG ATGCGTGAAT ACACCAGGCT
CATTTTATTG CCAGTGCAGT CCTGGGTTTC AATTGGCAGC AAACAACTAT ACCTGCGTAG
(SEQ ID No. 59)

Start intron 7

Gtaagccttttgagaacttgctgatttctgtcttcacaaaaggaaaccccatgccatgttgcagtatttccagt
tttctatgttcttgagtaaatagtttccatcgacttcccttcagcaatcataaagytgcaggagaaccccatga
agcttgaattcggatccacgtgg (SEQ ID No. 60)

End intron 7

aaactttcccactgaaagtgcattcttgatttttacatgccttttytccccctttcagaaatgcaatgtcacgg
:cacaacaattaaaagattggcatgacatgggaaaaaa:tctagtgtrggaaaaatgccttttcaacaatattt
t:cagtgctttagaagcattgcaaaactccgtatgggttctcaaaggcttatgttataattgtaatggaattta
acagaacccatttaaaaaagttaataa:atagccacagataaatcttccagtaccagcattgcctgaagaagac
catatccagtataagttgtcttatawcaattatttatagaaattggcattttgtwtcttgaaccaacaaaagaa
aaatccgaat:mccggaaktgttatatttwttagaagcattaaattcctttgganagattnatcacacatcnna
ctaactgtcattcctagaaaaaatatttcggtatttccnaaagaagtatatgacagacgtttgtagttgttccc
acaaatatganaccnaaatggatgttctccagtgagcttctgcagggcaaataattcagctagggaattactca
cttgtcagcagatgacgtaggtacaaaagagtaaggatatgtttaaagtstaymtatatmtgtgtgtgtataya
tatacatatacaymwmymtatayatamrattttttcwag (SEQ ID No. 61)

Exon 8

ATATAAATGA ATGTGATGCC AGCAATCAAT GTGCTCAGCA GTGCTACAAC ATTCTTGGTT
CATTCATCTG TCAGTGCAAT CAAGGATATG AGCTAAGCAG TGACAGGCTC AACTGTGAAG
(SEQ ID No. 62)

FIG.5C

Start intron 8

gtaaaactcttcccagatagtatggaataaagtccaattcctgtgactgctgttgttttatctgattatgta
cccatttatgaaaacaaacgcttacccatgggaattctgttctaaaatccagttaatgtagttcagttgtta
cattgccttttagtgtgttaccaagaaaaaggaaaagaaataaaaataactgaaatattaggtgcaggctg
gctctaataattagaaagggtgctctagcatgttgcgtctcagtgtgttatccagtgaccaggtatgtcagc
acctcctgggagcttatgagaaatgcagaatctcaggctgcaacccagacctcctgagtctgaatctacatt
caacaaaaactgcaggtgactggtgtgcacatttaagtttcagaatagccaagtgcatgtcaaaacattaaa
ataaaaatcaggagatctggtttctggttctattcctgctactgtgtgacttgggcacatttcttgagttgc
ctgggtttcactttccacatgaacaagaggagggccatttaactagattcatgaccttcagggtccattgca
tgtgcacatttcgttatataattcaaaaggcattagacatcctgaggggatgccacagacacttgatgtcc
ctgacctcctcacggttcactcagctttacacaaagctcaaaccccaccgagagaggcctcacatcatgcca
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actattcagaaatgcaaagcattcttcccctgaaagtcagatcctgccaagctgtcattctggaagcttgca
caggttaggggacttggcattcaaagctcaaatgaacttggcttcaaagtcacccaatttctgagaagacaa
acatgaactctacatcctggatgggtctgcagagtccaaaatgaaggccgtcaaccacaagccaattcattc
agtagtgtagttaggtccaggattagccaaattgtcagcaatgattcagtaaaagtcatgataagaaaaact
ttttgtgctatgaagtcatagagggaaataagctgatattgttagaatttgccttttagctgcttataaagt
tttgtatttctatttcagaatttgcaatatttttactctctttagctcacctcaaaagtgtattactt
cctctggactgttgagcagaaca (SEQ ID No. 63)

End intron 8

Aaaaaaatatatatgtgtgtgtgtgtgtgtgtgtgtgtgtgtatattaaacccagncaacttaaaaaatgt
gcccaagtcacacagtcgcaggaataggacaanaagccagatctctttatatatatataggta,gatataattt
ttcctccttanaatataaataa,ttttaattatatataattattttaatatagatattttaaatcttataattt
atatatatatataatttatatatatatatatatccaaagtagtggtgcacaaacttttcaactctgtgtcctt
tctcttgtctaattcaacag (SEQ ID No. 64)

Exon 9

ACATTGATGA ATGCAGAACC TCAAGCTACC TGTGTCAATA TCAATGTGTC AATGAACCTG GGAAATTCTC
ATGTATGTGC CCCCAGGGAT ACCAAGTGGT GAGAAGTAGA ACATGTCAAG
(SEQ ID No. 65)

Start intron 9

gtaagtttatttttttttcatatgttaggtatttagttttagccaggaagagacaagaggaagttataggatt
ctcctatagactttcatttttcccactttcaatatacaatttaagctnttttttcccctgttcatcataaaata
tatacatctcataaagaggggattctatgctaangccgacnttttttcgtccttaaaagataaataatttttaata
aaatattgatatgtattctatgtaacctacatcatctnttgagatacatcttcaaatcatccactggaaaaga
ttcagttattaaaangtttcacctgtgagtttgagtttanagcataagctcaatatgggagttaaacatacctc
catccagtcttagccctctaaaacncanggattataaattgcgtaaaaatgtaggtgctgaaaaaagtcagcct
aatatgttgtaaaatatagttgaatattttagagaaaactactagccccaaaatagctaatgaccttgggtcca
gtttcaaaataaacattcagatgatcttcacacctatacgtaagkggaagaggcagctccccacaatggtatga
tttcagagtttctcaggaagatctaaaaaaaaaaaggaccctacctccaatgttgcatgtagttgaaaattttc
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ttacncaaagcaccaaaagaatgaaaaaatgaccaancttgacanaaaacctgtttgaatcccagctccactgt
nttcagtctgcncaatnttgaacaaattatcaaactactntgagcctcagnttcctcatttggaaaagggagtt
gggggaatttaggggaatancatncntaaaaatantttgtaaactataaagcttgtncaggtcaaggggttttt
atnaaatttac (SEQ ID No. 66)

FIG.5D

End of intron 9

Agcctcttccttaacttcctcttttttccttacagtcctaaaattgctatgctctatgaggtggaacacttcata
gtttcacttcctgtgctgtgcttcctctggacagtataatccactcccagcatgcttcagcttactgaaaccag
atttctagcctttacctttctcccaagttcctgaaagagatgataagctgccctccatagtttatgcttcctga
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gagatgcctcttccttccttcagccttccctaatccccctggcaaaattacccaactctgctccacatgcccca
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gtccctgatgacaagaactggtgtttaacttttatataggcagagtcagtggttaacattggggtttgaattca
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gtctgtcctgaattcaatacaatgaacacccagtcttgtgtctaaaagcaggttgaacacagtccagatgctct
cacacctccttccttgcaaacagaatctgccagttatatgatttaattagatcagttcattagtttagttagta
aactctttgaccctacatctctacag (SEQ ID No. 67)

Exon 10

ATATAAATGA GTGTGAGACC ACAAATGAAT GCCGGGAGGA TGAAATGTGT TGGAATTATC
ATGGCGGCTT CCGTTGTTAT CCACGAAATC CTTGTCAAGA TCCCTACATT CTAACACCAG AGAA
(SEQ ID No. 68)

intron 10

gtaagaaaaa tcagaacttt tgaaagtgag gattttctgg tcttaccaagccaaactgct gaatactttt
gtttgtctct gcag
(SEQ ID No. 69)

Exon 11

CCGATGTGTT TGCCCAGTCT CAAATGCCAT GTGCCGAGAA CTGCCCCAGT CAATAGTCTA
CAAATACATG AGCATCCGAT CTGATAGGTC TGTGCCATCA GACATCTTCC AGATACAGGC
CACAACTATT TATGCCAACA CCATCAATAC TTTTCGGATT AAATCTGGAA ATGAAAATGG
AGAGTTCTAC CTACGA (SEQ ID No. 70)

Start intron 11

gtaagtatcc tgaaggcagc cttaactatt gagaaagatg ggagtttgttgttgttgttg ttgttgttgt
tgttgtgtgg tatccacatg tggaaggaaagcaaacattt aaaagtgtcttnatgtgtaggcattgtgtaag
gccttccagctacattatttcatttattcctcttggtaacactgccagatagatattaatattcatctccatttt
ttacagaggagaaaagtgagatgcagaaagattaagtagcatccctgaaatcactcaaatattaagtttggcaga
ctctgatagagttgtgtgtgaccacgaaaatacaagcctcccatccccccgcctctgccccacccaacatacc
cccaagtaggtatcactaatcattgatggttaattaattatacatagacatacatataattcaaacccaaaataa
ttcctggagctcctaaagagtttttcagacatcatgaattcatcattgttacattcacaagacagtttgtgttca
caccgaaactaaaacctataagtatgtgagaagtgaccccacctccccgcacagtatgtgtcaagtagttgtacc
ttcttgccaacttctgggctggcagtatggagtcatctccctatctttcattgcctgtgtgaaatctactttctg
aattctgccatttccctcttcacactgtctcctgggttatctttgcttcctcacatccctatctctcttcctata
aactggctcccgtcacttccatgatcccttcagtggcttctgagctggtctccctgaccccaaagcctcagcctt
ccagtctccctacaaaatctcagcaagttcatttaaggttaaaatttggacatattttaaatacggctcaccac
ttcatgtgaaaatgatggcaccctaccaagcagtttgcagagttaccggtaactgtttcatgctaatgatgttay
tcatccagttac SEQ ID No. 71)

End intron 11

Tccctttttttttttcyttctaaaaaggnaaccnatggcccaagnttgnaaaaanaaaaagggccnctttgnttt
ccaggtttaaaaatttccnattttcccctwaagtttagktttggaaaggcccccacttcnccannaaaaggaa
aaaaaatgnttacmaanagggggggattcaaaacnaaaaactttttttaaaaaaaaaaaaaaagcaagtccttgaa
acttggagctaatgactgtattagacaagggataagagccaagaagagttgaaaccaagaagggaccaagtagt
ggctcttttataccaccttcaaaattctccccctaattcttataggaggtatactaacaaagcatagaaactcc
aatccaagaaaattattctcttcctttctctatttttcttttatttag (SEQ ID No. 72)

FIG.5E

Exon 12

CAAACAAGTC CTGTAAGTGC AATGCTTGTG CTCGTGAAGT CATTATCAGG ACCAAGAGAA
CATATCGTGG ACCTGGAGAT GCTGACAGTC AGCAGTATAG GGACCTTCCG CACAAGCTCT
GTGTTAAGAT TGACAATAAT AGTGGGGCCA TTTTCATTTT AG (SEQ ID No. 73)

3' utr

tcttttctaa gagtcaacca caggcattta agtcagccaa agaatattgt taccttaaag
cactatttta tttatagata tatctagtgc atctacatct ctatactgta cactcaccca
taacaaacaa ttacaccatg gtataaagtg ggcatttaat atgtaaagat tcaaagtttg
tctttattac tatatgtaaattagacatta atccactaaa ctggtcttct tcaagagagc
taagtataca ctatctggtg aaacttggat tctttcctat aaaagtggga ccaagcaatg
atgatcttct gtggtgctta aggaaactta ctagagctcc actaacagtc tcataaggag
gcagccatca taaccattga atagcatgca agggtaagaatgagttttta actgctttgt
aagaaaatgg aaaaggtcaa taaagatatatttctttaga aaatggggat ctgccatatt
tgtgttggtt tttattttcatatccagcct aaaggtggtt gtttattata tagtaataaa
tcattgctgtacaacatgct ggtttctgta gggtattttt aattttgtca gaaattttagattgtgaata
ttttgtaaaa aacagtaagc aaaattttcc agaattcccaaaatgaacca gataccccct agaaaattat
actattgaga aatctatggggaggatatga gaaaataaat tccttctaaa ccacattgga
actgacctgaagaagcaaac tcggaaaata taataacatc cctgaattca ggcattcacaagatgcagaa
caaaatggat aaaaggtatt tcactggaga agttttaatttctaagtaaa atttaaatcc taacacttca
ctaatttata actaaaatttctcatcttcg tacttgatgc tcacagagga agaaaatgat
gatggtttttattcctggca tccagagtga cagtgaactt aagcaaatta ccctcctacccaattctatg
gaatatttta tacgtctcct tgtttaaaat ctgactgctttactttgatg tatcatattt ttaaataaaa
ataaatattc ctttagaagatcactctaaaa
(SEQ ID No. 74)

FIG.5F

M LKALFLTMLTL ALVKSQDTE ETITYTQCTD GYEWDPVRQQ
CKDIDECDIV PDACKGGMKC VNHYGGYLCL PKTAQIIVNN EQPQQETQPA
EGTSGATTGV VAASSMATSG VLPGGGFVAS AAAVAGPEMQ TGRNNFVIRR
NPADPQRIPS NPSHRIQCAA GYEQSEHNVC QDIDECTAGT HNCRADQVCI
NLRGSFACQC PPGYQKRGEQ CVDIDECTIP PYCHQRCVNT PGSFYCQCSP
GFQLAANNYT CVDINECDAS NQCAQQCYNI LGSFICQCNQ GYELSSDRLN
CEDIDECRTS SYLCQYQCVN EPGKFSCMCP QGYQVVRSRT CQDINECETT
NECREDEMCW NYHGGFRCYP RNPCQDPYIL TPENRCVCPV SNAMCRELPQ
SIVYKYMSIR SDRSVPSDIF QIQATTIYAN TINTFRIKSG NENGEFYLRQ
TSPVSAMLVL VKSLSGPREH IVDLEMLTVS SIGTFRTSSV LRLTIIVGPF
SF (SEQ ID NO.: 2)

FIG.6